

GenCore version 5.1.4\_p5\_4578  
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OM protein - protein search, using sw model

Run on: May 19, 2003, 16:43:42 ; Search time 19.1281 Seconds  
(without alignments)  
553.754 Million cell updates/sec

Title: US-09-625-573-4  
Perfect score: 1900  
Sequence: 1 MLSTSRFRINTNESGEEV.....DGVSTNTPTSGEQEVSAGL 360

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues  
Total number of hits satisfying chosen parameters: 262574

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Issued Patents AA: \*  
1: /cgn2.6/prodata/1/iaa/5A\_COMB.pep.\*  
2: /cgn2.6/prodata/1/iaa/5B\_COMB.pep.\*  
3: /cgn2.6/prodata/1/iaa/6A\_COMB.pep.\*  
4: /cgn2.6/prodata/1/iaa/6B\_COMB.pep.\*  
5: /cgn2.6/prodata/1/iaa/PCTUS\_COMB.pep.\*  
6: /cgn2.6/prodata/1/iaa/backfiles1.pep.\*

Pred. No. is the number of results predicted by chance to have a  
score greater than or equal to the score of the result being printed,  
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	1900	100.0	360	1	US-08-450-393A-4
2	1900	100.0	360	4	US-08-446-669-4
3	1900	100.0	360	4	US-09-045-583-50
4	1900	100.0	360	4	US-09-534-185-50
5	1900	100.0	360	5	PCT-US95-00476-4
6	1873	98.6	360	4	US-08-833-752-7
7	1849	97.3	360	4	US-09-045-583-51
8	1849	97.3	360	4	US-09-534-185-51
9	1838	96.7	347	1	US-08-461-244-3
10	1651.5	86.9	374	1	US-08-450-393A-2
11	1651.5	86.9	374	4	US-08-446-669-2
12	1651.5	86.9	374	5	PCT-US95-00476-2
13	1368.5	82.6	344	3	US-08-466-343D-9
14	1386	72.9	354	4	US-08-724-984A-2
15	1371	72.2	352	4	US-09-517-605-5
16	1370	72.1	352	4	US-09-045-583-52
17	1370	72.1	352	4	US-09-534-185-52
18	1364	71.8	352	4	US-09-087-232A-13
19	1364	71.8	352	4	US-08-861-105-14
20	1364	71.8	352	4	US-08-575-967A-2
21	1364	71.8	352	4	US-08-833-752-5
22	1355	71.3	352	3	US-08-466-343D-2
23	1036	54.5	355	1	US-08-012-988A-2
24	1036	54.5	355	1	US-08-450-393A-5
25	1036	54.5	355	4	US-08-446-669-5
26	1036	54.5	355	4	US-09-239-938-1
27	1036	54.5	355	5	PCT-US95-00476-5

28	1008	53.1	355	4	US-08-833-752-9	Sequence 9, Appli
29	981	51.6	355	4	US-09-045-583-53	Sequence 53, Appl
30	981	51.6	355	4	US-09-534-185-53	Sequence 53, Appl
31	947	49.8	355	4	US-08-575-967A-4	Sequence 4, Appli
32	947	49.8	355	4	US-08-847-296B-1	Sequence 1, Appli
33	947	49.8	355	4	US-09-045-583-54	Sequence 54, Appl
34	947	49.8	355	4	US-09-534-185-54	Sequence 54, Appl
35	917	48.3	355	4	US-08-833-752-8	Sequence 8, Appli
36	862.5	45.4	360	4	US-08-875-573-20	Sequence 20, Appl
37	862.5	45.4	360	4	US-09-232-878-2	Sequence 2, Appli
38	862.5	45.4	360	4	US-09-045-583-55	Sequence 55, Appl
39	862.5	45.4	360	4	US-09-534-185-55	Sequence 10, Appl
40	826.5	43.5	360	4	US-08-833-752-10	Sequence 2, Appli
41	751	39.5	355	1	US-08-461-244-2	Sequence 56, Appl
42	751	39.5	355	4	US-09-045-583-56	Sequence 56, Appl
43	751	39.5	355	4	US-09-534-185-56	Sequence 56, Appl
44	708	37.3	355	1	US-08-153-848-28	Sequence 28, Appl
45	708	37.3	355	1	US-08-153-848-32	Sequence 32, Appl

ALIGNMENTS

RESULT 1  
US-08-450-393A-4  
: Sequence 4, Application US/08450393A  
: Patent No. 5707815  
: GENERAL INFORMATION:  
: APPLICANT: Charo, Israel  
: TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMOATTRACTANT  
: TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMOATTRACTANT  
: NUMBER OF SEQUENCES: 14  
: CORRESPONDENCE ADDRESS:  
: ADDRESSEE: Cooley Godward Castro Huddleson & Tatum  
: STREET: 5 Palo Alto Square  
: CITY: Palo Alto  
: STATE: California  
: COUNTRY: USA  
: ZIP: 94306-2155  
: COMPUTER READABLE FORM:  
: MEDIUM TYPE: Floppy disk  
: OPERATING SYSTEM: IBM PC compatible  
: SOFTWARE: Patent Release #1.0, Version #1.25  
: CURRENT APPLICATION DATA:  
: APPLICATION NUMBER: US/08/450.393A  
: FILING DATE: May 25, 1995  
: CLASSIFICATION: 424  
: ATTORNEY/AGENT INFORMATION:  
: NAME: Cseri, Luann  
: REGISTRATION NUMBER: 31,822  
: REFERENCE/DOCKET NUMBER: UCAL-237/020US  
: TELECOMMUNICATION INFORMATION:  
: TELEPHONE: 415-843-5165  
: TELEFAX: 415-8857-0663  
: TELEX: 380816COOLEYPA  
: INFORMATION FOR SEQ ID NO: 4:  
: SEQUENCE CHARACTERISTICS:  
: LENGTH: 360 amino acids  
: TYPE: amino acid  
: TOPOLOGY: linear  
: MOLECULE TYPE: protein  
US-08-450-393A-4

Query Match 100.0%; Score 1900; DB 1; Length 360;  
Best Local Similarity 100.0%; Pred.No. 3.le-151;  
Matches 360; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MLSTSRFRINTNESGEEVTTFFDYDYGAPCHKFDVKQIGALLPPLYSLVFIQFVGN 60  
DB 1 MLSTSRFRINTNESGEEVTTFFDYDYGAPCHKFDVKQIGALLPPLYSLVFIQFVGN 60

QY 61 MLVVLILINCKKLCCLTDIYLLNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
DB 61 MLVVLILINCKKLCCLTDIYLLNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
QY 121 HIGYFGGIFIIILLTDRYLAIYHVAFAKARTVTFGVVTSVITLWAVFASVPGIIFTK 180  
DB 121 HIGYFGGIFIIILLTDRYLAIYHVAFAKARTVTFGVVTSVITLWAVFASVPGIIFTK 180  
QY 181 COKEDSVYVCGPYFPRGWNHFTIMRNILGLVLP LLIMVICYSIGILKTLRCRNEKKRHR 240  
DB 181 COKEDSVYVCGPYFPRGWNHFTIMRNILGLVLP LLIMVICYSIGILKTLRCRNEKKRHR 240  
QY 241 AVRVITIMIVFLFWTPYNNIVILLNTFQEFGLSNCESTSQLDQATQVTTGLMTHCCI 300  
DB 241 AVRVITIMIVFLFWTPYNNIVILLNTFQEFGLSNCESTSQLDQATQVTTGLMTHCCI 300  
QY 301 NPIIYAFVGEKFRRLYSVFFRKHTKRFCKQCPVYFRETVDGVTSTNTPSTGEQEVSA 360  
DB 301 NPIIYAFVGEKFRRLYSVFFRKHTKRFCKQCPVYFRETVDGVTSTNTPSTGEQEVSA 360

## RESULT 2

US-08-446-669-4  
; Sequence 4, Application US/08446669  
; Patent No. 6132987  
; GENERAL INFORMATION:  
; APPLICANT: Charo, Israel  
; APPLICANT: Coughlin, Shaun  
; TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMAOTTRACTANT  
; TITLE OF INVENTION: PROTEIN RECEPTORS  
; NUMBER OF SEQUENCES: 14  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Cooley Edward Castro Huddleson & Tatum  
; STREET: 5 Palo Alto Square  
; CITY: Palo Alto  
; STATE: California  
; COUNTRY: USA  
; ZIP: 94306-2155  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patent In Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/446,669  
; FILING DATE: May 25, 1995  
; CLASSIFICATION: 435  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Neeley, Richard  
; REGISTRATION NUMBER: 30,092  
; REFERENCE/DOCKET NUMBER: UCAL-237/01US  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 415-843-5000  
; TELEFAX: 415-857-0663  
; TELEX: 380816COOLEYPA  
; INFORMATION FOR SEQ ID NO: 4:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 360 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
US-08-446-669-4

Query Match 100.0%; Score 1900; DB 4; Length 360;  
Best Local Similarity 100.0%; Pred. No. 3.1e-151;  
Matches 360; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MLSTSRSRFIRNTNESGEVTTFFDYDYGAPCHKFDVKQIGAQLLPPLYSLVIFGFVGN 60  
DB 1 MLSTSRSRFIRNTNESGEVTTFFDYDYGAPCHKFDVKQIGAQLLPPLYSLVIFGFVGN 60  
QY 61 MLVVLILINCKKLCCLTDIYLLNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
DB 61 MLVVLILINCKKLCCLTDIYLLNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120

DB 61 MLVVLILINCKKLCCLTDIYLLNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
QY 121 HIGYFGGIFIIILLTDRYLAIYHVAFAKARTVTFGVVTSVITLWAVFASVPGIIFTK 180  
DB 121 HIGYFGGIFIIILLTDRYLAIYHVAFAKARTVTFGVVTSVITLWAVFASVPGIIFTK 180  
QY 181 COKEDSVYVCGPYFPRGWNHFTIMRNILGLVLP LLIMVICYSIGILKTLRCRNEKKRHR 240  
DB 181 COKEDSVYVCGPYFPRGWNHFTIMRNILGLVLP LLIMVICYSIGILKTLRCRNEKKRHR 240  
QY 241 AVRVITIMIVFLFWTPYNNIVILLNTFQEFGLSNCESTSQLDQATQVTTGLMTHCCI 300  
DB 241 AVRVITIMIVFLFWTPYNNIVILLNTFQEFGLSNCESTSQLDQATQVTTGLMTHCCI 300  
QY 301 NPIIYAFVGEKFRRLYSVFFRKHTKRFCKQCPVYFRETVDGVTSTNTPSTGEQEVSA 360  
DB 301 NPIIYAFVGEKFRRLYSVFFRKHTKRFCKQCPVYFRETVDGVTSTNTPSTGEQEVSA 360

## RESULT 3

US-09-045-583-50  
; Sequence 50, Application US/09045583  
; Patent No. 6287805  
; GENERAL INFORMATION:  
; APPLICANT: Graham, Gerard J. et al.  
; TITLE OF INVENTION: NO. 6287805el Molecules of the G Protein-Coupled  
; NUMBER OF SEQUENCES: 56  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: LAHIVE & COCKFIELD, LLP  
; STREET: 28 State Street  
; CITY: Boston  
; STATE: Massachusetts  
; COUNTRY: USA  
; ZIP: 02109  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patent In Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/09/045,583  
; FILING DATE: 20-MAR-98  
; CLASSIFICATION: 435  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER:  
; FILING DATE:  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Mandragoras, Amy E.  
; REGISTRATION NUMBER: 36,207  
; REFERENCE/DOCKET NUMBER: MNI-044  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (617)227-7400  
; TELEFAX: (617)742-4214  
; INFORMATION FOR SEQ ID NO: 50:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 360 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; FRAGMENT TYPE: internal  
US-09-045-583-50

Query Match 100.0%; Score 1900; DB 4; Length 360;  
Best Local Similarity 100.0%; Pred. No. 3.1e-151;  
Matches 360; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MLSTSRSRFIRNTNESGEVTTFFDYDYGAPCHKFDVKQIGAQLLPPLYSLVIFGFVGN 60  
DB 1 MLSTSRSRFIRNTNESGEVTTFFDYDYGAPCHKFDVKQIGAQLLPPLYSLVIFGFVGN 60  
QY 61 MLVVLILINCKKLCCLTDIYLLNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120  
DB 61 MLVVLILINCKKLCCLTDIYLLNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120

QY 121 HIGYFGGIFPIILLTIDRYLAIVHAVFALKARTVTFGVVTSVITLWVAVFASVPGIIFTK 180  
DB 121 HIGYFGGIFPIILLTIDRYLAIVHAVFALKARTVTFGVVTSVITLWVAVFASVPGIIFTK 180  
QY 181 COKEDSVYVCGPYFPRGNNFHTIMRNILGLVLPPLIMVTCYSGILKTLRCRNEKKRHR 240  
DB 181 COKEDSVYVCGPYFPRGNNFHTIMRNILGLVLPPLIMVTCYSGILKTLRCRNEKKRHR 240  
QY 241 AVRVFTIMIVYFLWTPYNNIVILLNTFQEFFGLSNCESSTQDQATQVTTGLMTHCCI 300  
DB 241 AVRVFTIMIVYFLWTPYNNIVILLNTFQEFFGLSNCESSTQDQATQVTTGLMTHCCI 300  
QY 301 NPIIYAFVGEKFRYLSVFFRKHITKRCQKCPVYFRETVDGVTSTNTPSTGEOEVSAGL 360  
DB 301 NPIIYAFVGEKFRYLSVFFRKHITKRCQKCPVYFRETVDGVTSTNTPSTGEOEVSAGL 360

RESULT 4  
US-09-534-185-50  
Sequence 50, Application US/09534185  
Patent No. 6403767  
GENERAL INFORMATION:  
APPLICANT: Graham, Gerard J. et al.  
TITLE OF INVENTION: No. 6403767el Molecules of the G Protein-Coupled  
Heptahelical Receptor Superfamily and Uses  
thereof  
NUMBER OF SEQUENCES: 56  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LAHIVE & COCKFIELD, LLP  
STREET: 28 State Street  
CITY: Boston  
STATE: Massachusetts  
COUNTRY: USA  
ZIP: 02109  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/534,185  
FILING DATE: 24-Mar-2000  
CLASSIFICATION: <Unknown>  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 09/045,583  
FILING DATE: <Unknown>  
ATTORNEY/AGENT INFORMATION:  
NAME: Mandragouras, Amy E.  
REGISTRATION NUMBER: 36,207  
REFERENCE/DOCKET NUMBER: MNI-044  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (617)227-7400  
TELEFAX: (617)742-4214  
INFORMATION FOR SEQ ID NO: 50:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 360 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FRAGMENT TYPE: internal  
SEQUENCE DESCRIPTION: SEQ ID NO: 50:  
US-09-534-185-50

Query Match 100.0%; Score 1900; DB 4; Length 360;  
Best Local Similarity 100.0%; Pred. No. 3.le-151;  
Matches 360; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MLSTSRSRFIRNTNESGEEVTFDDYDYGAPCHKFDVKQIGAOQLLPPLYSLVFIKFCVGN 60  
DB 1 MLSTSRSRFIRNTNESGEEVTFDDYDYGAPCHKFDVKQIGAOQLLPPLYSLVFIKFCVGN 60  
QY 61 MLVVLILINCKKLCITDIYLLNLAIISDLLFLITPLWAHSAANEWFNGNAMCKLFTGLY 120

DB 61 MLVVLILINCKKLCITDIYLLNLAIISDLLFLITPLWAHSAANEWFNGNAMCKLFTGLY 120  
QY 121 HIGYFGGIFPIILLTIDRYLAIVHAVFALKARTVTFGVVTSVITLWVAVFASVPGIIFTK 180  
DB 121 HIGYFGGIFPIILLTIDRYLAIVHAVFALKARTVTFGVVTSVITLWVAVFASVPGIIFTK 180  
QY 181 COKEDSVYVCGPYFPRGNNFHTIMRNILGLVLPPLIMVTCYSGILKTLRCRNEKKRHR 240  
DB 181 COKEDSVYVCGPYFPRGNNFHTIMRNILGLVLPPLIMVTCYSGILKTLRCRNEKKRHR 240  
QY 241 AVRVFTIMIVYFLWTPYNNIVILLNTFQEFFGLSNCESSTQDQATQVTTGLMTHCCI 300  
DB 241 AVRVFTIMIVYFLWTPYNNIVILLNTFQEFFGLSNCESSTQDQATQVTTGLMTHCCI 300  
QY 301 NPIIYAFVGEKFRYLSVFFRKHITKRCQKCPVYFRETVDGVTSTNTPSTGEOEVSAGL 360  
DB 301 NPIIYAFVGEKFRYLSVFFRKHITKRCQKCPVYFRETVDGVTSTNTPSTGEOEVSAGL 360

RESULT 5  
PCT-US95-00476-4  
Sequence 4, Application PC/TUS9500476  
GENERAL INFORMATION:  
APPLICANT: The Regents of the University of California  
TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMAATTRACTANT  
TITLE OF INVENTION: PROTEIN RECEPTORS  
NUMBER OF SEQUENCES: 14  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Robbins, Berliner & Carson  
STREET: 201 N. Figueroa Street, 5th Floor  
CITY: Los Angeles  
STATE: California  
COUNTRY: USA  
ZIP: 90012-2628  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: PCT/US95/00476  
FILING DATE:  
CLASSIFICATION:  
ATTORNEY/AGENT INFORMATION:  
NAME: Berliner, Robert  
REGISTRATION NUMBER: 20,121  
REFERENCE/DOCKET NUMBER: 5555-291  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 310-977-1001  
TELEFAX: 310-977-1003  
TELEX:  
INFORMATION FOR SEQ ID NO: 4:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 360 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
PCT-US95-00476-4

Query Match 100.0%; Score 1900; DB 5; Length 360;  
Best Local Similarity 100.0%; Pred. No. 3.le-151;  
Matches 360; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MLSTSRSRFIRNTNESGEEVTFDDYDYGAPCHKFDVKQIGAOQLLPPLYSLVFIKFCVGN 60  
DB 1 MLSTSRSRFIRNTNESGEEVTFDDYDYGAPCHKFDVKQIGAOQLLPPLYSLVFIKFCVGN 60  
QY 61 MLVVLILINCKKLCITDIYLLNLAIISDLLFLITPLWAHSAANEWFNGNAMCKLFTGLY 120  
DB 61 MLVVLILINCKKLCITDIYLLNLAIISDLLFLITPLWAHSAANEWFNGNAMCKLFTGLY 120  
QY 121 HIGYFGGIFPIILLTIDRYLAIVHAVFALKARTVTFGVVTSVITLWVAVFASVPGIIFTK 180

Db 121 HIGYGGIIFILLTIDRYLAIVHAVFALKARTVTFGVVTSITWLVAVFASVPGIIFTK 180  
QY 181 COKEDSVYVCGPYFPRGWNFNHTIMRNILGLVPLLLIMVICYSGLIKTLRCRNEKKRHR 240  
Db 181 COKEDSVYVCGPYFPRGWNFNHTIMRNILGLVPLLLIMVICYSGLIKTLRCRNEKKRHR 240  
QY 241 AVRVIETIMIVFLFWTPYINIVILLNTFQEFFGLSNCESTSQLDOAQVTTGLMTHCCI 300  
Db 241 AVRVIETIMIVFLFWTPYINIVILLNTFQEFFGLSNCESTSQLDOAQVTTGLMTHCCI 300  
QY 301 NPIIYAFVGEKFRYLSVFFRKHITKRFCKQCPVFEYRETVDGVTSTNTPSTGEOEVSAGL 360  
Db 301 NPIIYAFVGEKFRYLSVFFRKHITKRFCKQCPVFEYRETVDGVTSTNTPSTGEOEVSAGL 360

## RESULT 6

US-08-833-752-7

Sequence 7, Application: US/08833752

Patent No. 6448375

## GENERAL INFORMATION:

APPLICANT: SAMSON, MICHEL  
APPLICANT: PARMENTIER, MARC  
APPLICANT: VASSART, GILBERT  
APPLICANT: LIBERT, FREDERICK  
TITLE OF INVENTION: ACTIVE AND INACTIVE CC-CHEMOKINES RECEPTOR  
TITLE OF INVENTION: AND NUCLEIC ACID MOLECULES ENCODING SAID RECEPTOR  
NUMBER OF SEQUENCES: 17  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Knobbe, Martens, Olson & Bear  
STREET: 620 Newport Center Drive 16th Floor  
CITY: Newport Beach  
STATE: CA  
COUNTRY: U.S.A.  
ZIP: 92660

## COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent In Release #1.0, Version #1.25 (EPO)  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/833,752  
FILING DATE: 9-APR-1997  
CLASSIFICATION: 536  
ATTORNEY/AGENT INFORMATION:  
NAME: Altman, Daniel E

REGISTRATION NUMBER: 34,115

REFERENCE/DOCKET NUMBER:

INFORMATION FOR SEQ ID NO: 7:

SEQUENCE CHARACTERISTICS:

LENGTH: 360 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: No. 6448375e

US-08-833-752-7

Query Match 98.6%; Score 1873; DB 4; Length 360;

Best Local Similarity 98.3%; Pred. No. 5.6e-149;

Matches 354; Conservative 2; Mismatches 4; Indels 0; Gaps 0;

QY 1 MLSTSRFRIRNTNSESSEVTTFFDYDYGAPCHKFDVKQIGALLPPLYSLVFIQFVGN 60  
Db 1 MLSTSRFRIRNTNSESSEVTTFFDYDYGAPCHKFDVKQIGALLPPLYSLVFIQFVGN 60  
QY 61 MLVVLILNCKKLKCLTDIYLLNLAIISDLLFLITPLWAHSAANEVWFGNAMCKLFTGLY 120  
Db 61 MLVVLILNCKKLKCLTDIYLLNLAIISDLLFLITPLWAHSAANEVWFGNAMCKLFTGLY 120  
QY 121 HIGYGGIIFILLTIDRYLAIVHAVFALKARTVTFGVVTSITWLVAVFASVPGIIFTK 180  
Db 121 HIGYGGIIFILLTIDRYLAIVHAVFALKARTVTFGVVTSITWLVAVFASVPGIIFTK 180

QY 181 COKEDSVYVCGPYFPRGWNFNHTIMRNILGLVPLLLIMVICYSGLIKTLRCRNEKKRHR 240  
Db 181 COKEDSVYVCGPYFPRGWNFNHTIMRNILGLVPLLLIMVICYSGLIKTLRCRNEKKRHR 240  
QY 241 AVRVIETIMIVFLFWTPYINIVILLNTFQEFFGLSNCESTSQLDOAQVTTGLMTHCCI 300  
Db 241 AVRVIETIMIVFLFWTPYINIVILLNTFQEFFGLSNCESTSQLDOAQVTTGLMTHCCI 300  
QY 301 NPIIYAFVGEKFRYLSVFFRKHITKRFCKQCPVFEYRETVDGVTSTNTPSTGEOEVSAGL 360  
Db 301 NPIIYAFVGEKFRYLSVFFRKHITKRFCKQCPVFEYRETVDGVTSTNTPSTGEOEVSAGL 360

## RESULT 7

US-09-045-583-51

Sequence 51, Application US/09045583

Patent No. 6287805

## GENERAL INFORMATION:

APPLICANT: Graham, Gerard J. et al.  
TITLE OF INVENTION: NO. 6287805el Molecules of the G Protein-Coupled  
NUMBER OF SEQUENCES: 56  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LAHIVE & COCKFIELD, LLP  
STREET: 28 State Street  
CITY: Boston  
STATE: Massachusetts  
COUNTRY: USA  
ZIP: 02109

## COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent In Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/045,583  
FILING DATE: 20-MAR-98  
CLASSIFICATION: 435  
PRIOR APPLICATION NUMBER:  
APPLICATION NUMBER:  
FILING DATE:  
ATTORNEY/AGENT INFORMATION:  
NAME: Mandragouras, Amy E.

REGISTRATION NUMBER: 36,207

REFERENCE/DOCKET NUMBER: MNI-044

TELEPHONE: (617)227-7400

TELEFAX: (617)742-4214

INFORMATION FOR SEQ ID NO: 51:

SEQUENCE CHARACTERISTICS:

LENGTH: 360 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: peptide

FRAGMENT TYPE: internal

US-09-045-583-51

Query Match 97.3%; Score 1849; DB 4; Length 360;

Best Local Similarity 97.2%; Pred. No. 5.6e-147;

Matches 350; Conservative 5; Mismatches 5; Indels 0; Gaps 0;

QY 1 MLSTSRFRIRNTNSESSEVTTFFDYDYGAPCHKFDVKQIGALLPPLYSLVFIQFVGN 60  
Db 1 MLSTSRFRIRNTNSESSEVTTFFDYDYGAPCHKFDVKQIGALLPPLYSLVFIQFVGN 60  
QY 61 MLVVLILNCKKLKCLTDIYLLNLAIISDLLFLITPLWAHSAANEVWFGNAMCKLFTGLY 120  
Db 61 MLVVLILNCKKLKCLTDIYLLNLAIISDLLFLITPLWAHSAANEVWFGNAMCKLFTGLY 120  
QY 121 HIGYGGIIFILLTIDRYLAIVHAVFALKARTVTFGVVTSITWLVAVFASVPGIIFTK 180  
Db 121 HIGYGGIIFILLTIDRYLAIVHAVFALKARTVTFGVVTSITWLVAVFASVPGIIFTK 180  
QY 181 COKEDSVYVCGPYFPRGWNFNHTIMRNILGLVPLLLIMVICYSGLIKTLRCRNEKKRHR 240

Db 181 COEDSVVYCGPYPRGWNNEHTIMRNILGLVLPILLIMVICYSGLTKLLRCRNEKKRHR 240  
QY 241 AVRVIETIMIVYFWPTPNIVILLNTFOEFGLSNCESTSLDQATQVTTGLMTHCCI 300  
Db 241 AVRVIETIMIVYFWPTPNIVILLNTFOEFGLSNCESTSLDQATQVTTGLMTHCCI 300  
QY 301 NPIIYAFVGEKFRRLYSVFFRKHTIKRCKQCPVYFRETVDGVTSTNTPSTGEQVSAGL 360  
Db 301 NPIIYAFVGEKFRRLYSVFFRKHTIKRCKQCPVYFRETVDGVTSTNTPSTGEQVSAGL 360

## RESULT 8

US-09-534-185-51  
Sequence-51, Application US/09534185  
Patent No. 6403767  
GENERAL INFORMATION:  
APPLICANT: Graham, Gerard J. et al.  
TITLE OF INVENTION: No. 6403767el Molecules of the G Protein-Coupled  
Heptahelical Receptor Superfamily and Uses  
Therefor

NUMBER OF SEQUENCES: 56  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: LAHIVE & COCKFIELD, LLP  
STREET: 28 State Street  
CITY: Boston  
STATE: Massachusetts  
COUNTRY: USA  
ZIP: 02109

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/534,185  
FILING DATE: 24-Mar-2000  
CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 09/045,583  
FILING DATE: <Unknown>  
ATTORNEY/AGENT INFORMATION:  
NAME: Mandragouras, Amy E.  
REGISTRATION NUMBER: 36,207  
REFERENCE/DOCKET NUMBER: MNI-044  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (617)227-7400  
TELEFAX: (617)742-4214

INFORMATION FOR SEQ ID NO: 51:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 360 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FRAGMENT TYPE: internal  
SEQUENCE DESCRIPTION: SEQ ID NO: 51:

US-09-534-185-51

Query Match 97.38; Score 1849; DB 4; Length 360;  
Best Local Similarity 97.24; Pred. No. 5.6e-147;  
Matches 350; Conservative 5; Mismatches 5; Indels 0; Gaps 0;

QY 1 MLSTSRSRFIRNTNCSGEVTTFFDYDYGAPCHKFDVKQIGAOQLLPPLYSLVFIFGFGVGN 60  
Db 1 MLSTSRSRFIRNTNCSGEVTTFFDYDYGAPCHKFDVKQIGAOQLLPPLYSLVFIFGFGVGN 60  
QY 61 MLVVLILINCKKLCLETDIYLLNLAIISDLLFLITPLWAHSAANEWVFGNAMCKLFTGLY 120  
Db 61 MLVVLILINCKKLSLTDIYLLNLAIISDLLFLITPLWAHSAANEWVFGNAMCKLFTGLY 120  
QY 121 HIGYFGGIFFTLLIDRYLAIVHAVFALKARTVTFGVVTSVITWLVAVFASVPGIIFTK 180  
Db 121 HIGYFGGIFFTLLIDRYLAIVHAVFALKARTVTFGVVTSVITWLVAVFASVPGIIFTK 180

QY 181 COKEDSVVYCGPYPRGWNNEHTIMRNILGLVLPILLIMVICYSGLTKLLRCRNEKKRHR 240  
Db 181 COEDSVVYCGPYPRGWNNEHTIMRNILGLVLPILLIMVICYSGLTKLLRCRNEKKRHR 240  
QY 241 AVRVIETIMIVYFWPTPNIVILLNTFOEFGLSNCESTSLDQATQVTTGLMTHCCI 300  
Db 241 AVRVIETIMIVYFWPTPNIVILLNTFOEFGLSNCESTSLDQATQVTTGLMTHCCI 300  
QY 301 NPIIYAFVGEKFRRLYSVFFRKHTIKRCKQCPVYFRETVDGVTSTNTPSTGEQVSAGL 360  
Db 301 NPIIYAFVGEKFRRLYSVFFRKHTIKRCKQCPVYFRETVDGVTSTNTPSTGEQVSAGL 360

## RESULT 9

US-08-461-244-3  
Sequence 37, Application-US/08461244  
Patent No. 5776729  
GENERAL INFORMATION:  
APPLICANT: Soppet, Daniel R.  
APPLICANT: Yi, Li  
APPLICANT: Ruben, Steven M.  
APPLICANT: Rosen, Craig A.  
TITLE OF INVENTION: HUMAN G-PROTEIN RECEPTOR HGBER32  
NUMBER OF SEQUENCES: 7  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN, CECCHI,  
ADDRESSEE: STUART & OLSTEIN  
STREET: 6 Becker Farm Road  
CITY: Roseland  
STATE: New Jersey  
COUNTRY: USA  
ZIP: 07068

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/461,244  
FILING DATE: 05-JUN-1995  
CLASSIFICATION: 536

ATTORNEY/AGENT INFORMATION:  
NAME: Ferraro, Gregory D.  
REGISTRATION NUMBER: 36,134  
REFERENCE/DOCKET NUMBER: 325800-445  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 201-994-1700  
TELEFAX: 201-994-1744

INFORMATION FOR SEQ ID NO: 3:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 347 amino acids  
TYPE: amino acid  
STRANDEDNESS:  
TOPOLOGY: linear  
MOLECULE TYPE: protein

US-08-461-244-3

Query Match 96.7%; Score 1838; DB 1; Length 347;  
Best Local Similarity 100.0%; Pred. No. 4.5e-146;  
Matches 347; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 14 NESGEVTTFFDYDYGAPCHKFDVKQIGAOQLLPPLYSLVFIFGFGVGNMLVLLINCKKL 73  
Db 1 NESGEVTTFFDYDYGAPCHKFDVKQIGAOQLLPPLYSLVFIFGFGVGNMLVLLINCKKL 60  
QY 74 KCLTDIYLLNLAIISDLLFLITPLWAHSAANEWVFGNAMCKLFTGLYHIGYFGGIFFT 133  
Db 61 KCLTDIYLLNLAIISDLLFLITPLWAHSAANEWVFGNAMCKLFTGLYHIGYFGGIFFT 120  
QY 134 LTIDRYLAIVHAVFALKARTVTFGVVTSVITWLVAVFASVPGIIFTKCKEDSVVYCGPY 193  
Db 121 LTIDRYLAIVHAVFALKARTVTFGVVTSVITWLVAVFASVPGIIFTKCKEDSVVYCGPY 180

QY 194 FPRGWNFFHIMRNILGLVLPPLIMVICYSGILKTLRCRNEKKRRHRAVRVIFTIMIYVF 253  
|||||  
Db 181 FPRGWNFFHIMRNILGLVLPPLIMVICYSGILKTLRCRNEKKRRHRAVRVIFTIMIYVF 240  
|||||  
QY 254 LEWTPYNIIVLLNTFOEFFGLSNCESTSQLDOATQVTTGLMTHCCINPIIYAVGKFR 313  
|||||  
Db 241 LEWTPYNIIVLLNTFOEFFGLSNCESTSQLDOATQVTTGLMTHCCINPIIYAVGKFR 300  
|||||  
QY 314 RYLSVFRKHITKRKCKQCPVFXREYVDGVTSTNTPTSGEQEVSAGL 360  
|||||  
Db 301 RYLSVFRKHITKRKCKQCPVFXREYVDGVTSTNTPTSGEQEVSAGL 347  
|||||

## RESULT 10

US-08-450-393A-2

; Sequence 2, Application US/08450393A

; Patent No. 5707815

; GENERAL INFORMATION:

; APPLICANT: Charo, Israel

; APPLICANT: Coughlin, Shaun

; TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMOATTRACTANT

; TITLE OF INVENTION: PROTEIN RECEPTORS

; NUMBER OF SEQUENCES: 14

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Cooley Godward Castro Huddleson &amp; Tatum

; STREET: 5 Palo Alto Square

; CITY: Palo Alto

; STATE: California

; COUNTRY: USA

; ZIP: 94306-2155

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: Patent In Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/450,393A

; FILING DATE: May 25, 1995

; CLASSIFICATION: 424

; ATTORNEY/AGENT INFORMATION:

; NAME: Cserit, Luann

; REGISTRATION NUMBER: 31,822

; REFERENCE/DOCKET NUMBER: UCAL-237/02US

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: 415-843-5165

; TELEFAX: 415-8857-0663

; TELEX: 380816CooLeYPA

; INFORMATION FOR SEQ ID NO: 2:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 374 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: protein

US-08-450-393A-2

Query Match 86.9%; Score 1651.5; DB 1; Length 374;

Best Local Similarity 95.5%; Pred. No. 1.8e-130;

Matches 319; Conservative 3; Mismatches 5; Indels 7; Gaps 3;

QY 1 MLSTSRFRIRNTNBSGEEVTTFFDYDGAPCHKFDVKQIGAQLLPPLSLVFIQFVGN 60  
|||||  
Db 1 MLSTSRFRIRNTNBSGEEVTTFFDYDGAPCHKFDVKQIGAQLLPPLSLVFIQFVGN 60  
|||||  
QY 61 MLVVLILNCKKLCTDIYLLNLALISDLFLITPLWAHSAANEVFGNAMCKLFTGLY 120  
|||||  
Db 61 MLVVLILNCKKLCTDIYLLNLALISDLFLITPLWAHSAANEVFGNAMCKLFTGLY 120  
|||||  
QY 121 HIGYFGGFFIILLTDRLYLAVHAVFALKARTVTEGVVTSVITLWVAVFASVPGIIFTK 180  
|||||  
Db 121 HIGYFGGFFIILLTDRLYLAVHAVFALKARTVTEGVVTSVITLWVAVFASVPGIIFTK 180  
|||||  
QY 181 CQKEDSVYVCGPYFPRGWNFFHIMRNILGLVLPPLIMVICYSGILKTLRCRNEKKRRH 240  
|||||

Db 181 CQKEDSVYVCGPYFPRGWNFFHIMRNILGLVLPPLIMVICYSGILKTLRCRNEKKRRH 240  
|||||  
QY 241 AVRVIPTIMIVFLFWTPYPYNIIVLLNTFOEFFGLSNCESTSQLDOATQVTTGLMTHCCI 300  
|||||  
Db 241 AVRVIPTIMIVFLFWTPYPYNIIVLLNTFOEFFGLSNCESTSQLDOATQVTTGLMTHCCI 300  
|||||  
QY 301 NPIIYAFVGEKFRRLVSVFRRKHITKRKCKQCPV 334  
|||||  
Db 301 NPIIYAFVGEKFRRLVSVFRRKHITKRKCKQCPV 327  
|||||

## RESULT 11

US-08-446-669-2

; Sequence 2, Application US/08446669

; Patent No. 6132987

; GENERAL INFORMATION:

; APPLICANT: Charo, Israel

; APPLICANT: Coughlin, Shaun

; TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMOATTRACTANT

; TITLE OF INVENTION: PROTEIN RECEPTORS

; NUMBER OF SEQUENCES: 14

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Cooley Godward Castro Huddleson &amp; Tatum

; STREET: 5 Palo Alto Square

; CITY: Palo Alto

; STATE: California

; COUNTRY: USA

; ZIP: 94306-2155

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: Patent In Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/446,669

; FILING DATE: May 25, 1995

; CLASSIFICATION: 435

; ATTORNEY/AGENT INFORMATION:

; NAME: Neeley, Richard

; REGISTRATION NUMBER: 30,092

; REFERENCE/DOCKET NUMBER: UCAL-237/01US

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: 415-843-5000

; TELEFAX: 415-857-0663

; TELEX: 380816CooLeYPA

; INFORMATION FOR SEQ ID NO: 2:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 374 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: protein

US-08-446-669-2

Query Match 86.9%; Score 1651.5; DB 4; Length 374;

Best Local Similarity 95.5%; Pred. No. 1.8e-130;

Matches 319; Conservative 3; Mismatches 5; Indels 7; Gaps 3;

QY 1 MLSTSRFRIRNTNBSGEEVTTFFDYDGAPCHKFDVKQIGAQLLPPLSLVFIQFVGN 60  
|||||  
Db 1 MLSTSRFRIRNTNBSGEEVTTFFDYDGAPCHKFDVKQIGAQLLPPLSLVFIQFVGN 60  
|||||  
QY 61 MLVVLILNCKKLCTDIYLLNLALISDLFLITPLWAHSAANEVFGNAMCKLFTGLY 120  
|||||  
Db 61 MLVVLILNCKKLCTDIYLLNLALISDLFLITPLWAHSAANEVFGNAMCKLFTGLY 120  
|||||  
QY 121 HIGYFGGFFIILLTDRLYLAVHAVFALKARTVTEGVVTSVITLWVAVFASVPGIIFTK 180  
|||||  
Db 121 HIGYFGGFFIILLTDRLYLAVHAVFALKARTVTEGVVTSVITLWVAVFASVPGIIFTK 180  
|||||  
QY 181 CQKEDSVYVCGPYFPRGWNFFHIMRNILGLVLPPLIMVICYSGILKTLRCRNEKKRRH 240  
|||||  
Db 181 CQKEDSVYVCGPYFPRGWNFFHIMRNILGLVLPPLIMVICYSGILKTLRCRNEKKRRH 240  
|||||

QY 241 AVRVIITMIVYFLWTPYINIVILLNTFOEFFGLSNCESTSQLDOATQVETLGMTHCCI 300  
Db 241 AVRVIITMIVYFLWTPYINIVILLNTFOEFFGLSNCESTSQLDOATQVETLGMTHCCI 300  
QY 301 NPIIYAFVGEKFRYLSVFFKHKHITKRFCKQCPV 334  
Db 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327

## RESULT 12

PCT-US95-00476-2  
Sequence 2, Application PC/TUS9500476  
GENERAL INFORMATION:

APPLICANT: The Regents of the University of California

TITLE OF INVENTION: MAMMALIAN MONOCYTE CHEMOTRACTANT

TITLE OF INVENTION: PROTEIN RECEPTORS

NUMBER OF SEQUENCES: 14

CORRESPONDENCE ADDRESS:

ADDRESSEE: Robbins, Berliner & Carson

STREET: 201 N. Figueroa Street, 5th Floor

CITY: Los Angeles

STATE: California

COUNTRY: USA

ZIP: 90012-2628

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patent In Release #1.0, Version #1.25

CURRENT APPLICATION NUMBER: PCT/US95/00476

FILING DATE:

CLASSIFICATION:

ATTORNEY/AGENT INFORMATION:

NAME: Berliner, Robert

REGISTRATION NUMBER: 20,121

REFERENCE/DOCKET NUMBER: 5555-291

TELECOMMUNICATION INFORMATION:

TELEPHONE: 310-977-1001

TELEFAX: 310-977-1003

TELEX:

INFORMATION FOR SEQ ID NO: 2:

SEQUENCE CHARACTERISTICS:

LENGTH: 374 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

PCT-US95-00476-2

Query Match 86.9%; Score 1651.5; DB 5; Length 374;  
Best Local Similarity 95.5%; Pred. No. 1.8e-130;  
Matches 319; Conservative 3; Mismatches 5; Indels 7; Gaps 3;

QY 1 MLSTSRSPRIRTNNSGEEVTFDDYDYGAPCHKFDVKQIGAQLLPPLYSLSVIFGVGN 60  
Db 1 MLSTSRSPRIRTNNSGEEVTFDDYDYGAPCHKFDVKQIGAQLLPPLYSLSVIFGVGN 60  
QY 61 MLVLLINCKLKCLTDIYLLNLAISDLLFLITPLWAHSAANWVFGNCKLFTGLY 120  
Db 61 MLVLLINCKLKCLTDIYLLNLAISDLLFLITPLWAHSAANWVFGNCKLFTGLY 120  
QY 121 HIGYEGGIFFIILLTDIYLAIVHAFVFAKARTVTFGVVTSVITLWVAFASVPGIIFTK 180  
Db 121 HIGYEGGIFFIILLTDIYLAIVHAFVFAKARTVTFGVVTSVITLWVAFASVPGIIFTK 180  
QY 181 COKEDSVYVCGPYFPGWNEHTIMRNILGLVPLIMVICYSGILKTLRCRNEKRRH 240  
Db 181 COKEDSVYVCGPYFPGWNEHTIMRNILGLVPLIMVICYSGILKTLRCRNEKRRH 240  
QY 241 AVRVIITMIVYFLWTPYINIVILLNTFOEFFGLSNCESTSQLDOATQVETLGMTHCCI 300  
Db 241 AVRVIITMIVYFLWTPYINIVILLNTFOEFFGLSNCESTSQLDOATQVETLGMTHCCI 300

QY 301 NPIIYAFVGEKFRYLSVFFKHKHITKRFCKQCPV 334  
Db 301 NPIIYAFVGEKFR---SLF---HIALG-CRIAPL 327

## RESULT 13

US-08-466-343D-9

Sequence 9, Application-US/08466343D

Patent No. 6025154

GENERAL INFORMATION:

APPLICANT: LI, YI

TITLE OF INVENTION: POLYNUCLEOTIDES ENCODING HUMAN G-PROTEIN

TITLE OF INVENTION: CHEMOKINE RECEPTOR HDGNR10 (AS AMENDED)

NUMBER OF SEQUENCES: 9

CORRESPONDENCE ADDRESS:

ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C.

STREET: 1100 NEW YORK AVE., NW, SUITE 600

CITY: WASHINGTON

STATE: DC

COUNTRY: USA

ZIP: 20005

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patent In Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/466,343D

FILING DATE: 06-JUN-1995

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: STEFFE, ERIC K.

REGISTRATION NUMBER: 36,688

REFERENCE/DOCKET NUMBER: 1488.1150000/EKS/KLM

TELECOMMUNICATION INFORMATION:

TELEPHONE: (202) 371-2600

TELEFAX: (202) 371-2540

INFORMATION FOR SEQ ID NO: 9:

SEQUENCE CHARACTERISTICS:

LENGTH: 344 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: protein

US-08-466-343D-9

Query Match 82.6%; Score 1568.5; DB 3; Length 344;  
Best Local Similarity 95.3%; Pred. No. 1.4e-123;  
Matches 302; Conservative 3; Mismatches 5; Indels 7; Gaps 3;

QY 18 BEVTTFDDYDYGAPCHKFDVKQIGAQLLPPLYSLSVIFGVGNMVLINCKKLC 77  
Db 1 BEVTTFDDYDYGAPCHKFDVKQIGAQLLPPLYSLSVIFGVGNMVLINCKKLC 60  
QY 78 DIYLLNLAISDLLFLITPLWAHSAANWVFGNCKLFTGLYHIGYEGGIFILLTID 137  
Db 61 DIYLLNLAISDLLFLITPLWAHSAANWVFGNCKLFTGLYHIGYEGGIFILLTID 120  
QY 138 RYLAIVHAFVFAKARTVTFGVVTSVITLWVAFASVPGIIFTKCQEDSVYVCGPYFPRG 197  
Db 121 RYLAIVHAFVFAKARTVTFGVVTSVITLWVAFASVPGIIFTKCQEDSVYVCGPYFPRG 180  
QY 198 WNEHTIMRNILGLVPLIMVICYSGILKTLRCRNEKRRHRAVRVITMIVYFLFWT 257  
Db 181 WNEHTIMRNILGLVPLIMVICYSGILKTLRCRNEKRRHRAVRVITMIVYFLFWT 240  
QY 258 PYNIVILLNTFOEFFGLSNCESTSQLDOATQVETLGMTHCCINPIIYAFVGEKFRYLS 317  
Db 241 PYNIVILLNTFOEFFGLSNCESTSQLDOATQVETLGMTHCCINPIIYAFVGEKFRYLS 317  
QY 318 VFFRKHITKRFCKQCPV 334  
Db 318 VFFRKHITKRFCKQCPV 334

